

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0 Q9UL12	SARDH_HUMAN	Homo sapiens	Sarcosine dehydrogenase, mitochondrial	18.379595	NaN	T777	30379171	MASLSRALRVAAAHPRQSPTRGMGP CNLSSAAGPTAEKSVYPYQRTLKEGQ GTSVVAQGPSRPLPSTANVVVIGGGS LGCQTLYHLAKLGMMSGAVLLERERL TSGTTWHTAGLLWQLRPSDVEVELL AHTRRVVSRELEEEETGLHTGWIQNG GLFIASNRQRLDEYKRLMSLGKAYG VESHVLSPAETKTYPLMNVDDLYG TLYVPHDGTMDPAGTCTTLARAASA RGAQVIENCPTGIRVWTDDFGVRR VAGVETOHGSIQTPCVVNCAGVWAS AVGRMAGVKVPLVAMHHAYVVERI EGIQNMPNVRDHDASVYLRQLQGD LSVGGYEANPIFWEEVSDKFAFGLF DLDWEVFTQHIEGAINRVPVLEKTI KSTVCGPESFTPDHKPLMGEAPELR GFFLGC GFNSAGMMLGGGCGQELA HWIIGRPEKDMHGYDIRRFHSLT DHPRWIRERSHESYAKNYSVVFP EPLAGRNMRRDPLHEELLGQGCVF QERHCWERPCWFHPRGPAPVLEYD YYGAYGSRAHEDYAYRRLLADEYTF FPPHDTIKKECLACRGAADVFDMS YFGKFYLVGLDARKAADWLFADVS RPPGSTVYTCMLNHRGTESDLTVS RLAPSHQASPLAPAFEGDGYLAMG GAVAQHNWSHITTVLQDQKSQCQLI DSSDLGMISIQGPASRAILQEVLD DLSNEAFPSTHKLRAAGHLVRAM RLSFVGE LGWELHIPKASCVPVYRA VMAAGAKHGLINAGYRAIDSLSIEKG YRHWHADLRPDDSPLEAGLAFTCKL KSPVFLGREALEQQRAGLRRLV CFTMEDKVPMPGLEAIWRNGQVVG HVRRADFGFAIDKTIAYGYIHDPSGG PVSLDFVKSGDYALERMGVTYGAQA HLKSPFDPNNKRKVGII	True	False	2.432	1.446	5.0	1.066	0.694	1.154	1.268